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SEQUENCE LISTING
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GENERAL INFORMATION:

(i)

APPLICANT: PEREGRINO FERREIRA, Paulo;

5 GESSIEN KROON, Erna;

PIMENTA DOS REIS, Karlisson Jennner;

BIAS FORTES FERRAZ, Isabella;

CERQUEIRA LEITE, Romulo.

(ii)

TITLE OF INVENTION: Method and composition for the diagnosis of equine infectious anemia virus disease by using the recombinant capsid protein virus (p26)

(iii)

NUMBER OF SEQUENCES: 1

15 (iv)

CORRESPONDENCE ADDRESS:

(A)

ADDRESSEE: Universidade Federal de Minas Gerais - CTIT

(B)

20 STREET: Avenida Antônio Carlos, 6627 Bairro São Francisco

(C)

CITY: Belo Horizonte

(D)

STATE: Minas Gerais

25 (E)

COUNTRY: BRAZIL

(F)

ZIP: 31270-901

(v)

30 COMPUTER READABLE FORM:

(A)

MEDIUM TYPE: diskette – 3.50 inch, 1.44 Mb storage

(B)

COMPUTER: IBM compatible

(C)

5 OPERATING SYSTEM: Windows 98

(D)

SOFTWARE: Office premium

(vi)

CURRENT APPLICATION DATA:

10 (A)

APPLICATION NUMBER: U.S. 09/331.262

(B)

FILING DATE:

(C)

15 CLASSIFICATION: C12Q1/70

(vii)

PRIOR APPLICATION DATA

(A)

APPLICATION NUMBER: PI 9606273-8

20 (B)

FILING DATE: 18-DEC-1996

(2)

INFORMATION FOR SEQ ID N0:1:

(i)

25 SEQUENCE CHARACTERISTICS:

(A)

LENGHT: 252 amino acids

(B)

TYPE: amino acid

30 (D)

TOPOLOGY: linear

```
(ii)
    MOLECULE TYPE: protein
    (vi)
    ORIGINAL SOURCE
    (A)
    ORGANISM: equine infectious anemia virus
    (ix)
    FEATURE:
10
    (A)
    NAME: p26
    (x)
    PUBLICATION INFORMATION
    (A)
    AUTHORS:
15
   (B)
    TITLE: (
    C)
    JOURNAL:
    (D)
20
    VOLUME:
    (F)
    PAGES:
    (G)
    DATE:
25
    (xi)
    SEQUENCE DESCRIPTION: SEQ ID NO:1
    His His His His His Gly Ser Pro Gly Asn Pro Leu Thr Trp
                                                    15
30
                  5
                                   10
```

	Ser Lys Ala Leu Lys Lys Leu Glu Lys Val Thr Val Gln Gly Ser			
	20	25	30	
	Gin Lys Leu Thr Thr Gly Asn Cys Na Trp Ala Leu Ser Leu Val			
	35	40	45	
5	Asp Leu Phe His Asp Thr	Asn Phe Val Lys Glu Lys As	p Trp Gln	
	50	55	60	
	Leu Arg Asp Val Ile Pro Le	eu Leu Glu Asp Val Thr Gln	Thr Val	
	65	70	75	
	Ser Gly Gln Glu Arg Glu A	la Phe Glu Arg Thr Trp Trp	Ala Ile	
10	80	85	90	
	Ser Ala Val Lys Met Gly L	eu Gln lle Asn AsnVal Val A	sp Gly	
	95	100	105	
	Lys Ala Ser Phe Gln Leu L	_eu Arg Ala Lys Tyr Glu Lys	Lys Thr	
	110	115	120	
15	Ala Asn Lys Lys Gln Ser G	Glu Pro Ser Glu Glu Tyr Pro	lle Met	
	125	130	135	
	lle Asp Gly Ala Gly Asn Ar	rg Asn Phe Arg Pro Leu Thr	Pro Arg	
	140	145	150	
	Gly Tyr Thr Thr Trp Val As	snThr lle Gln Thr Asn Gly Le	u Leu	
20	155	160	165	
	Asn Glu Ala Ser Gln Asn I	Leu Phe Gly lle Leu Ser Val	Asp Cys	
	170	175	180	
	Thr Ser Glu Glu Met Asn	Ala Phe Leu Asp Val Val Pro	Gly Gln	
	185	190	195	
25	Ala Gly Gln Lys Gln Ile Le	u Leu Asp Ala lle Asp Lys II	e Ala	
	200	205	210	
	Asp Asp Trp Asp Asn Arg	His Pro Leu Pro Asn Ala Pr	o Leu Val	
٠.	215	220	225	
	Ala Pro Pro Gln Gly Pro II	e Pro Met Thr Ala Arg Phe I	le Arg	
30	230	235	240	
	Gly Leu Gly Vai Pro Arg Glu Arg Gln Met Glu Pro			
	245	250		

	Asn Cys Val Val Gln Ser Phe Gly Val Ile Gly Gln Ala His Leu.			
	260	265	270	
	Glu Leu Pro Arg Pro Asn Lys Arg Ile Arg Asn Gln. Ser Phe Asn			
	275	280	285	
5	Gln Tyr Asn Cys Ser lle Asn. Asn Lys Thr Glu Leu Glu Thr Trp			
	290	295	300	
	Lys Leu.Val Lys Thr Ser Gly Val Thr Pro Leu Pro. lle Ser Ser			
	305	310	315	
	Glu Ala Asn Thr Gly Leu			
10	320			